

# CONFIRMATION OF PHENCYCLIDINE BY GAS CHROMATOGRAPHY - MASS SPECTROMETRY

#### 2.1 METHOD

This test method may be used to confirm the presence of phencyclidine (PCP) in biological specimens. The target compound and phencyclidine-d<sub>5</sub> (PCP-d<sub>5</sub>) internal standard are isolated from biological matrices by solid phase extraction (SPE). The extracts are injected into a gas chromatograph (GC) coupled to a mass spectrometer (MS) detector equipped with an electron ionization source.

#### 2.2 SPECIMENS

The specimen volume is 1 mL. Specimens include, but are not limited to, whole blood, serum, plasma, urine, and tissue homogenate. Dilutions of specimens may be analyzed at the Forensic Scientist's discretion.

NOTE: Method validation established that matrix-matching of the full calibration curve and all positive control levels is not required for quantitation in liver (tissue) homogenate specimens (see 2.4.3.4). Matrix-matching of the full curve and all positive controls is required for quantitation in serum/plasma specimens (see 2.4.2 and 2.4.3).

### 2.3 REAGENTS, MATERIALS AND EQUIPMENT

#### 2.3.1 REAGENTS

NOTE: Organic solvents used are reagent grade.

- Acetic acid (glacial)
- 0.1M Acetic acid

Add 5.72 mL glacial acetic acid to 800 mL DI  $H_2O$ . Dilute to 1 L with DI  $H_2O$  and mix. Store the acid in a glass bottle at room temperature for up to 6 months.

- Acetonitrile (ACN)
- Ammonium hydroxide (NH<sub>4</sub>OH), concentrated
- Certified blank blood and/or other biological matrices
- Deionized water (DI H<sub>2</sub>O), laboratory general-use
- Elution solvent

To 20 mL isopropanol, add 2 mL concentrated ammonium hydroxide and mix. Add 78 mL methylene chloride and mix. Store the solvent in a glass flask/bottle at room temperature and use on date of preparation only.

- Ethyl acetate
- Iso-octane
- Isopropanol (IPA)
- Methanol (MeOH)



- Methylene chloride (dichloromethane, CH<sub>2</sub>Cl<sub>2</sub>)
- 0.1M Phosphate buffer (pH6):

Dissolve 1.7 g Na<sub>2</sub>HPO<sub>4</sub> and 12.14 g NaH<sub>2</sub>PO<sub>4</sub> • H<sub>2</sub>O in 800 mL DI H<sub>2</sub>O. Dilute to 1 L with DI H<sub>2</sub>O and mix. Check the pH and, if necessary, adjust to pH6  $\pm$  0.5 with concentrated NaOH or HCI. Store the buffer in a glass bottle at room temperature for up to one year.

- Sodium phosphate, dibasic anhydrous (Na<sub>2</sub>HPO<sub>4</sub>)
- Sodium phosphate, monobasic monohydrate (NaH₂PO₄ H₂O)

NOTE: Adjustments to final volumes of prepared reagents are permitted as long as the proportions are maintained.

#### 2.3.2 MATERIALS

- Disposable extraction tubes (16 x 100 mm recommended) and screw-cap or centrifuge tubes with closures
- Extraction column: United Chemical Technologies' Clean Screen SPE cartridge (CSDAU206, 200 mg/6 mL), or equivalent
- GC column (Agilent HP-5MS; 30 m x 0.250 mm i.d. x 0.250 µm film thickness, or equivalent)
- Glass autosampler vials with inserts and caps
- Laboratory glassware (graduated cylinders, flasks)

#### 2.3.3 EQUIPMENT

- Agilent GC (6890 or equivalent)
- Agilent MS (5973 or equivalent) with electron ionization source
- Calibrated, adjustable piston pipettes and verified, adjustable repeater-pipette with disposable pipette tips
- General-use equipment (centrifuge, evaporator, pH meter or paper, vacuum manifold, vortex mixer)

#### 2.4 STANDARDS, CALIBRATORS AND CONTROLS

#### 2.4.1 STANDARDS

Working standard: 10 ng/µL
Working control standard: 10 ng/µL
Working internal standard (PCP-d₅): 1 ng/µL

#### 2.4.2 CALIBRATORS

Calibrators are prepared in certified blank blood at the time of analysis, as detailed in 2.5 SAMPLE PREPARATION. Quantitation in serum/plasma specimens requires that a calibration curve be prepared in blank matrix. If testing only serum/plasma specimens, a whole blood calibration curve is not required.



#### 2.4.3 CONTROLS

- 2.4.3.1 At least one negative whole blood control and two positive whole blood controls are included in the batch, prepared as described in 2.5. For quantitative analysis of serum/plasma specimens only, whole blood controls are not required.
- 2.4.3.2 Controls (positive or negative) must make up at least 10% of the extracted batch (based on number of case specimen samples), with case specimens bracketed by positive controls.
- 2.4.3.3 For qualitative analysis of any alternate matrices, one negative control and one positive control must be included for each alternate matrix type tested in the batch.
- 2.4.3.4 For quantitative analysis of liver (tissue) homogenate specimens, matrixmatching of the full calibration curve and all positive controls is not required. One negative control and one positive control must be included in the batch. Positive controls in both whole blood and/or tissue homogenate serve to bracket tissue homogenate specimens.
- 2.4.3.5 For quantitative analysis of serum/plasma specimens, matrix-matching of the full calibration curve and all positive controls (to meet 10% and bracket specimens in that matrix) is required.

#### 2.5 SAMPLE PREPARATION

- 2.5.1 Label a clean extraction tube for each member of the test batch. (i.e., calibrator, control, case sample).
- 2.5.2 Add 4 mL of 0.1M phosphate buffer (pH6) into each tube.
- 2.5.3 Using a calibrated pipette, add 1 mL of certified blank whole blood into each of the calibrator tubes, positive control tubes, and negative control tube(s).
- 2.5.4 Prepare a 1:10 dilution of the working standard. (1 ng/µL)
  - a. Using a calibrated pipette, combine 0.1 mL of the working standard with 0.9 mL of ACN or MeOH in a labeled tube.
  - b. Cap and vortex mix. This dilution shall be disposed of after calibrator preparation.
- 2.5.5 Prepare a 1:100 dilution of the working standard.  $(0.1 \text{ ng/}\mu\text{L})$ 
  - c. Using a calibrated pipette, combine 0.1 mL of the 1:10 dilution with 0.9 mL of ACN or MeOH in a labeled tube.
  - d. Cap and vortex mix. This dilution shall be disposed of after calibrator preparation.
- 2.5.6 Using a calibrated pipette, spike the calibrators according to the following table, using the working standard and the prepared dilutions.



Calibrator Description	Volume (µL) Added	Standard Concentration	Dilution of WS (or WS)
Calibrator 1 – 10 ng/mL	100	0.1 ng/μL	1:100
Calibrator 2 – 25 ng/mL	25	1 ng/μL	1:10
Calibrator 3 – 50 ng/mL	50	1 ng/μL	1:10
Calibrator 4 – 100 ng/mL	100	1 ng/μL	1:10
Calibrator 5 - 500 ng/mL	50	10 ng/μL	WS
Calibrator 6 - 1000 ng/mL	100	10 ng/μL	WS

- 2.5.7 Prepare a 1:10 dilution of the control working standard. (1 ng/µL)
  - a. Using a calibrated pipette, combine 0.1 mL of the control working standard with 0.9 mL of ACN or MeOH in a labeled tube.
  - b. Cap and vortex mix. This dilution shall be disposed of after control preparation.
- 2.5.8 Using a calibrated pipette, spike the positive controls according to the following table, using the control working standard and prepared dilution.

Control	Volume (µL)	Standard	Dilution of
Description	Added	Concentration	QC (or QC)
Control 1 – 30 ng/mL	30	1 ng/μL	1:10
Control 2 - 250 ng/mL	25	10 ng/μL	QC

- 2.5.9 Using a calibrated pipette, sample 1 mL of each case sample into its respective tube.
- 2.5.10 Using a calibrated pipette or verified repeater-pipette, add 100  $\mu$ L of the working internal standard solution to each tube. Final concentration of the internal standard is 100 ng/mL.
- 2.5.11 Cap the tubes and briefly vortex mix. Centrifuge the tubes for 10 minutes at 3500 rpm (recommended for 16 x 100 mm tubes).
- 2.5.12 Place new, labeled SPE columns into the vacuum manifold.
- 2.5.13 Condition the SPE columns by passing each of the following solvents completely through under force of gravity.
  - a. 3 mL methanol
  - b. 3 mL DI H<sub>2</sub>O
  - c. 1 mL 0.1M phosphate buffer (pH6)

Do not let columns dry out between each conditioning step.

2.5.14 Transfer the contents of each tube to its respective SPE column and allow them to flow through under force of gravity. (Moderate, positive pressure or vacuum may



be applied if the flow is insufficient.)

- 2.5.15 Wash the SPE columns by passing each of the following solvents completely through under force of gravity. (Moderate, positive pressure or vacuum may be applied if the flow is insufficient.)
  - a. 3 mL DI H<sub>2</sub>O
  - b. 2 mL 0.1M acetic acid
  - c. 3 mL methanol
- 2.5.16 Dry the columns for 10 minutes under vacuum.
- 2.5.17 Place clean, labeled centrifuge or screw-cap tubes in the collection rack underneath their corresponding SPE columns.
- 2.5.18 Pass 3 mL elution solvent through each SPE column and collect the extracts.
- 2.5.19 Transfer the tubes to the evaporator and evaporate the extracts to dryness at 50°C.
- 2.5.20 Reconstitute the extracts by the addition of 50 µL ethyl acetate to each tube. Briefly vortex mix the tubes. If necessary, centrifuge the tubes for 2 minutes at 2000 rpm to collect the extracts at the bottom of the tubes.
- 2.5.21 Transfer the extracts to labeled glass autosampler vials with inserts and cap.

#### 2.6 INSTRUMENTAL PARAMETERS/DATA ANALYSIS

- Acquisition method PCP SIM (instrumental parameters in Appendix A)
- Calibration curve linear, 1/a weighting factor
- Updating calibrator (retention times ±2%, ion ratios ±20%) Cal 4
- Result comparisons –

Cal 1: truncated to one decimal place in units of ng/mL (acceptable range 7.5-12.5 ng/mL)

Cals 2-6, Ctls 1-2: truncated, whole integer values in units of ng/mL

#### 2.7 REPORTING

Results are truncated to two significant figures for reporting, in units of milligrams per liter (mg/L).

#### 2.8 METHOD PERFORMANCE

- Limit of detection: 1 ng/mL (0.001 mg/L)
- Lower limit of quantification: 10 ng/mL (0.010 mg/L)
- Dynamic range: 10 1000 ng/mL (0.010 1.0 mg/L)
- Upper limit of quantitation: 1000 ng/mL (1.0 mg/L)



## APPENDIX A INSTRUMENTAL PARAMETERS

## GAS CHROMATOGRAPH

Split/Splitless Inlet		
Mode	Split	
Temperature	250° C	
Split Ratio	2:1	
Gas Type	Helium	
Gas Saver	Off	
Gas Saver Flow	N/A	
Gas Saver Time	N/A	
Autosampler		
Injection Volume	2.0 μL	
Solvent Wash A	3 (Isooctane)	
Solvent Wash B	3 (Ethyl acetate)	
Sample Pumps	2	

Oven/Column		
Carrier Gas Mode	Constant Flow	
Carrier Gas Flow	1.0 mL/min	
Initial Temperature	150° C	
Initial Time	1.00 min	
Ramp Rate	15° C/min	
Final Temperature	290° C	
Final Time	0.67 min	
Transfer Line Temp	280 ° C	

### MASS SPECTROMETER

Solvent Delay	3.00 min	MS Quad Temperature	150° C
EM Offset	Set in Tune	MS Source Temperature	230° C
Resolution	Low	Dwell Time	50 msec
Signals	lons	Ion Ratios	
Phencyclidine	200, 243, 242	243/200, 242/200	
Phencyclidine-d <sub>5</sub>	205, 248	248/205	



## **LIST OF CHANGES**

Revision		
Date	Description	Page Number
09/01/11	Method approved by Washington State Toxicologist. See DRA dated 8/18/11. Method released for use in evidentiary testing on 09/01/11.	All
2/4/16	Added wording for adjustment of prepared volumes in 2.5.1.2, 2.5.1.12, 2.6.1.3 and 2.6.1.4 and clarification to 2.6.3.2.c for use of same CRM in preparation of working standard and working control standard. Added note regarding CRM expiration dates to 2.6.1.3 and 2.6.1.4. Edited 2.12.3 to reflect that only two significant figures are used for reporting and added "Printed Copies are Uncontrolled" to footer. Other minor edits throughout.	All
5/8/17	Wording added to 2.4.3 regarding dilution and standard volume testing. Specified use of calibrated pipettes for measurement of blank blood, specimens and standards throughout sample preparation in 2.7. Specified calibrator concentration criteria/ranges in 2.10.1.3. Edited 2.10.2.2.d to indicate all positive controls must pass for a target compound to report quantitative results. Other minor edits throughout.	1-8
8/8/19	Removed policy, purpose and principle sections, summarizing under new section METHOD. Added specific wording regarding matrix-matching in 2.2 SPECIMENS, 2.4.2 CALIBRATORS and 2.4.3 CONTROLS. Edited STANDARDS section - this information is now included in the revised Standard Solution Preparation procedure. Criteria for batch acceptance (calibrators, controls) and specimen acceptability criteria, and specific data analysis and reporting information are now included in the General Requirements for Chromatographic Test Method Batch Analysis and Acceptance. Formatting and minor edits throughout.	All